

Table 1

Fly primers for qRT-PCR

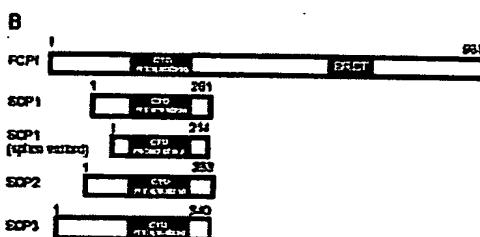
Name of gene	5' sequence	3' sequence	Accession no.
SCP1	5' atggcgaactatacgagtcgttc 3'	5' cttgtctgctgtggtaacatgg 3'	<u>CG5830</u>
GAPDH	5' atcaacgcacaacttcgagatcg 3'	5' gcggttggagtagccaaactcg 3'	<u>CG1205_5</u>
ribosomal protein S35	5' atgtcgcttgcaaaaactaagc 3'	5' ttataggatatctcgatttcggc 3'	<u>CG5497</u>
beta-actin	5' tgaagatccaccggagcgccgt 3'	5' gaccggactcgcatactcctgc 3'	<u>NM_079486</u>
Na Channel II	5' cagctggcggagtgacggcttc 3'	5' tgccgacgtcgccccatgttagac 3'	<u>CG9071</u>
synapsin	5' gagctgtcggtgagcttggcg 3'	5' cgctggattttggaaagaaggc 3'	<u>CG3985</u>
cholineAcetylTra nsferase	5' actggccattactacttgctc 3'	5' ccgtaaaaccgcgcgcattaaagt 3'	<u>CG3284_8</u>
ELAV	5' caacgaagccgagcgagccatcc 3'	5' tggcatggcacgaaatccgaatc 3'	<u>CG4396</u>
beta-tubulin	5' gcaacaactggccaagggttac 3'	5' ctggcatcgaaacatctgtgggtc 3'	<u>CG9277</u>
Neurofilament H	5' gcctccaagagcacgtacaaag 3'	5' cgatcagaagtggatcgccctta 3'	<u>CG7421</u>
peptidyl-glycine oxygenase	5' ctgcatttcaagtaccttgtgtc 3'	5' ccctggctgaagcagaacttcatg 3'	<u>CG3832</u>
myosin-light-chain-kinase	5' cttcgctgcacccatcagaaacgt 3'	5' tatggcataaaagggtggccattc 3'	<u>CG1915</u>
GCM	5' caacggaaactaacggccgtccgag 3'	5' gttctgccatgttggatctgc 3'	<u>CG1224_5</u>
nMDAR	5' ctcgcattttctctgggg 3'	5' cgtacatggatggacccctgg 3'	<u>CG1479_3</u>

Mouse Primers for RT-PCR

Name of gene	5' sequence	3' sequence	Accession no.
SCP1	5' cggccgttattactcagatcagcaagg 3'	5' gcagtgaacagcacacattcaaagg 3'	<u>AY028804</u>
GAPDH	5' tccaccaccctgtgtgtctgt 3'	5' accacatccatgcctac 3'	<u>NM_008084</u>
ngn1	5' catctctgtatctcgactgtccagg 3'	5' gggtcagagagtggatgcccac 3'	<u>NM_010896</u>
beta-tubulin	5' tgcctcaccatggggatctgttgc 3'	5' cttgaacagatccctggatggc 3'	<u>NM_023716</u>
stra13	5' ctgtggccatggggaaacagtggcttc 3'	5' agaagtccaggagcagctgggg 3'	<u>NM_016665</u>
GAD1	5' gcaaccgcaggcacgtttacggag 3'	5' agatgaccatccggaaagaagtggccctgt 3'	<u>NM_008077</u>
nrsf	5' ccattgcctgcgaaacctccccagg 3'	5' agccaaactcagctggactctcc 3'	<u>NM_011263</u>

Human Primers for ChIP assay

Name of gene	5' sequence	3' sequence	Accession no.
GAD1 promoter chr2q31	5' tgcgttatattatcctgcacgggg 3'	5' caccgggtcgagtcccgagggat 3'	<u>NT005403</u>
GAD1 3' gene chr2q31	5' ggagccatgcaggtaagggataa 3'	5' gggcttgcattttggagccacc 3'	<u>NT005403</u>
GRIN 2A promoter chr16	5' aactattctgggtactccttagac 3'	5' gctggaggaaatgcatttaatgc 3'	<u>NT010393</u>
SCN2 promoter chr.2q23	5' ctggataaggtaactgaagagtgg 3'	5' cagacgacaaggatcatgc 3'	<u>NT005403</u>



**FIG. 1**

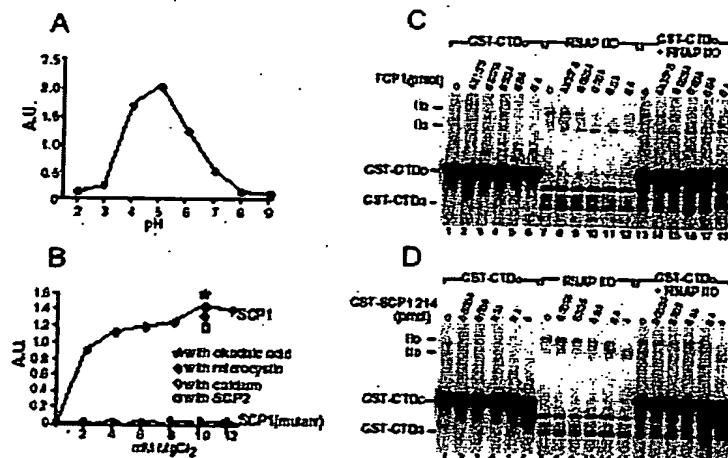
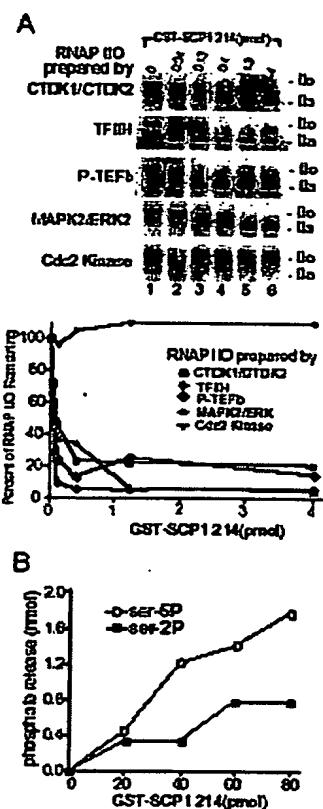


FIG. 2

**FIG. 3**

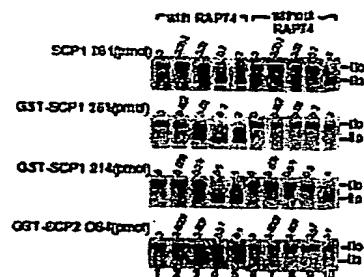


FIG. 4

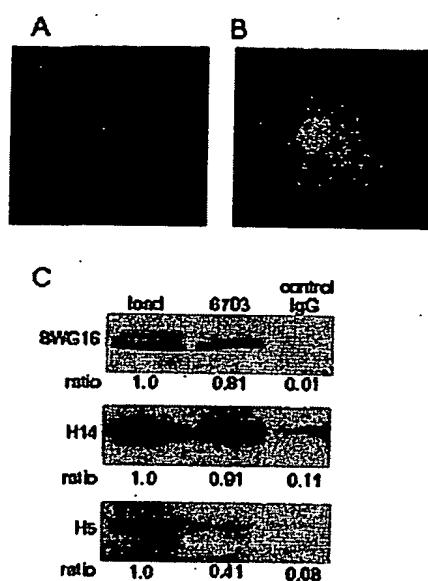
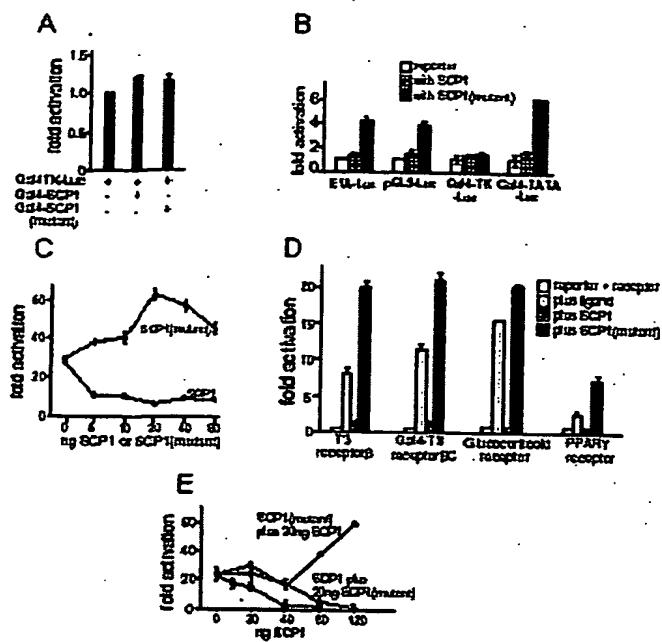


FIG. 5

**FIG. 6**

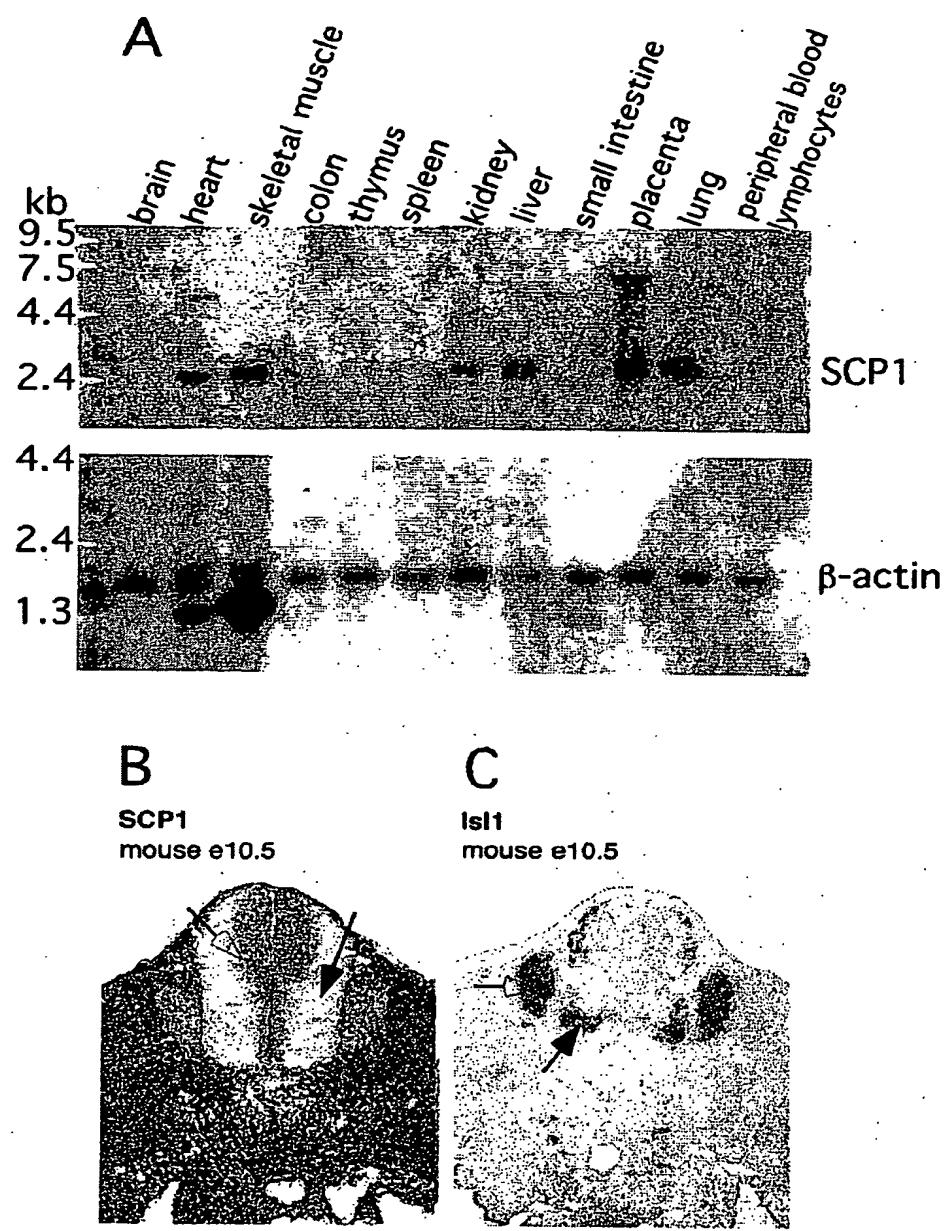


Figure 7

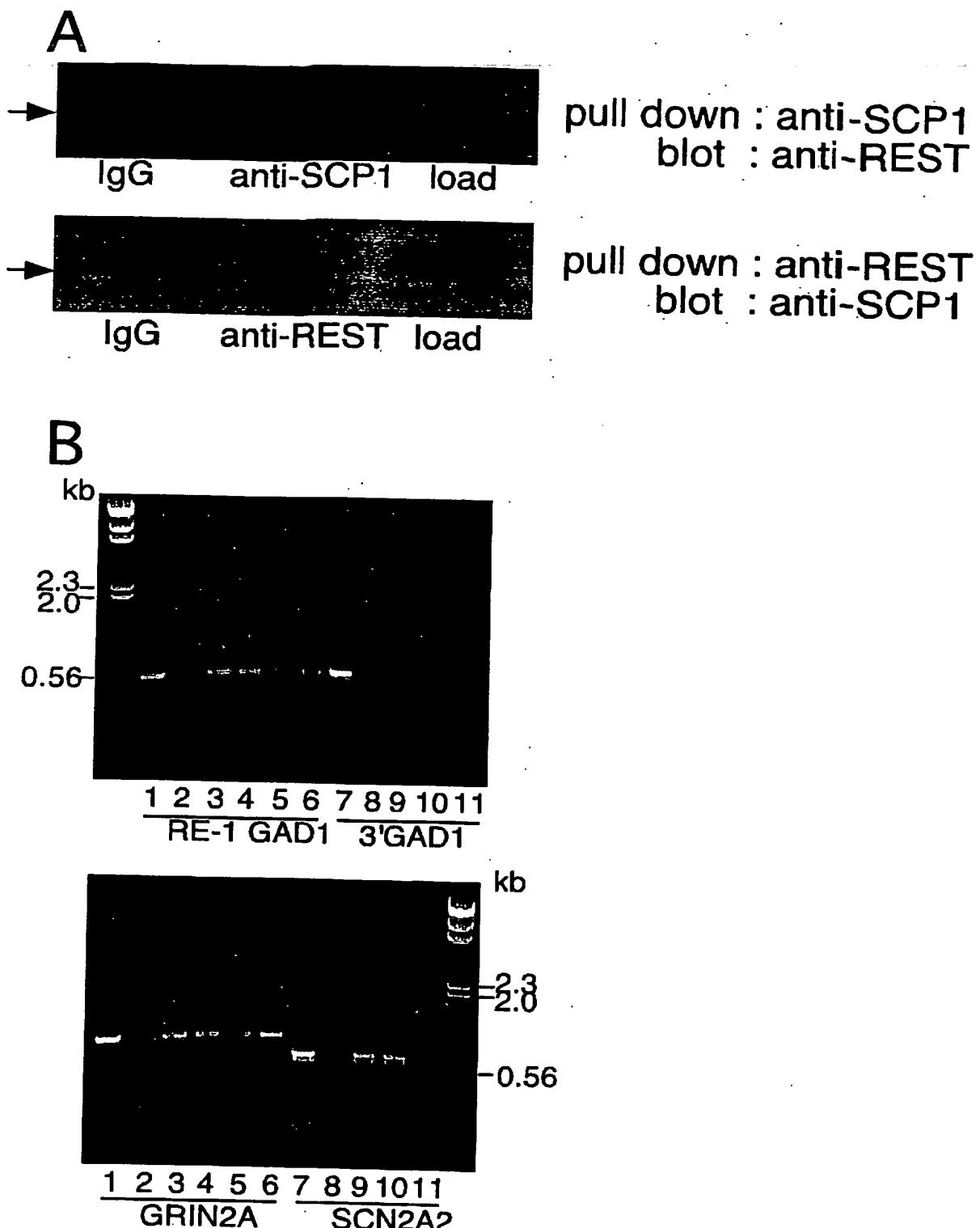
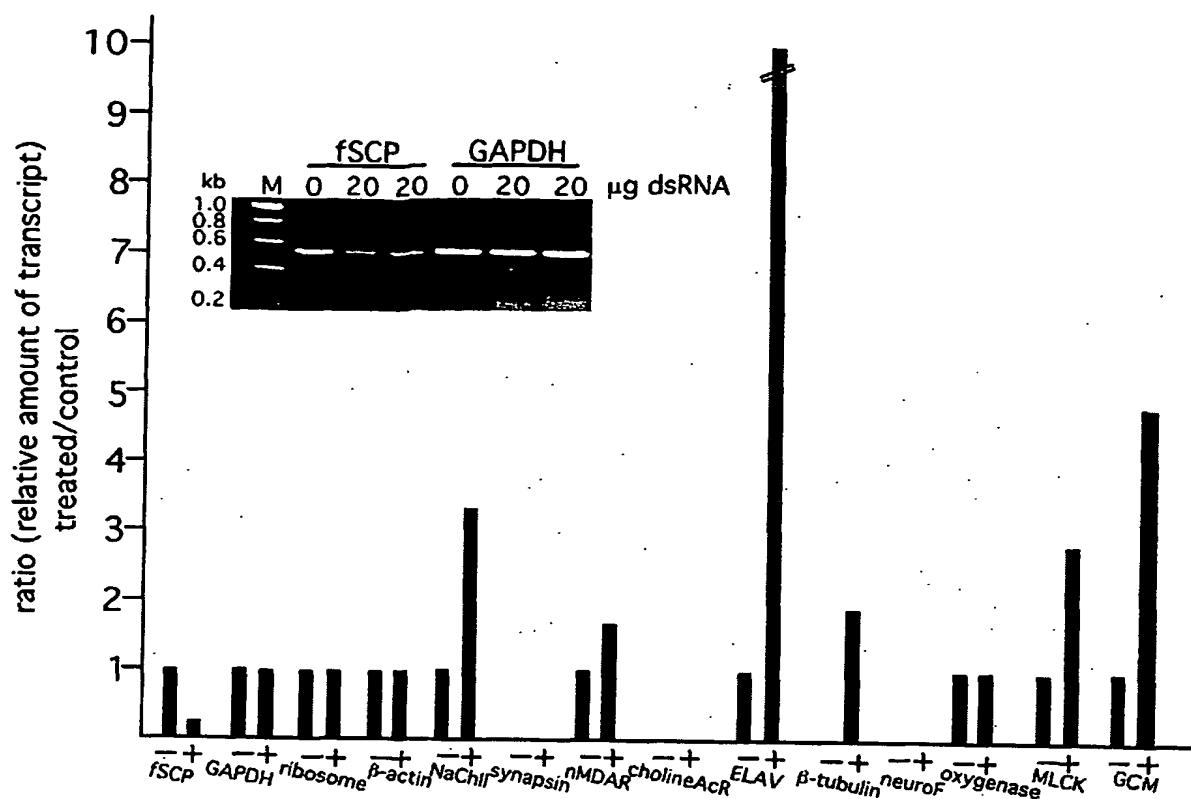


Figure 8



\*  $p = 0.001$  compared to wildtype  
\*\*  $p = 0.008$  compared to wildtype

Figure 9



gene		average Ct	std deviation
fSCP	control	21.1	0.18
	knockdown	23.3	0.22
GAPDH	control	17	0.1
	knockdown	17	0.19
ribosome	control	25	0.14
	knockdown	25	0.15
$\beta$ -actin	control	15	0.1
	knockdown	15	0.91
NaChII	control	25.1	0.22
	knockdown	23.8	0.2
synapsin	control	29.5	0.26
	knockdown	29.5	0.4
nMDAR	control	15.8	0.5
	knockdown	15.1	0.16
cholineAcR	control	29.4	0.76
	knockdown	31.2	0.96
ELAV	control	26.8	0.19
	knockdown	22	0.69
$\beta$ -tubulin	control	27.9	0.67
	knockdown	25.1	0.35
neuroF	control	30.2	0.83
	knockdown	30.1	0.55
oxygenase	control	23	0.46
	knockdown	23	2.16
MLCK	control	17.8	0.21
	knockdown	16.5	0.68
GCM	control	23	0.42
	knockdown	21	1.16

**Figure 10**